



PCT10

## RAW SEQUENCE LISTING

DATE: 08/06/2002

PATENT APPLICATION: US/10/069,111A

TIME: 16:29:01

Input Set : A:\p22033.ST25.txt

Output Set: N:\CRF3\08062002\J069111A.raw

**Does Not Comply  
Corrected Diskette Needed**

3 <110> APPLICANT: YANAGAWA, Hiroshi  
 4 NEMOTO, Naoto  
 W--> 5 <120> TITLE OF INVENTION: Method for Analyzing Interaction between Protein and Molecule  
 W--> 6 <130> FILE REFERENCE: P22033  
 W--> 7 <140> CURRENT APPLICATION NUMBER: 10/069,111A  
 8 <141> CURRENT FILING DATE: 2002-02-27  
 9 <150> PRIOR APPLICATION NUMBER: PCT/JP00/05920  
 10 <151> PRIOR FILING DATE: 2000-08-31  
 W--> 11 <160> NUMBER OF SEQ ID: 16  
 12 <170> SOFTWARE: PatentIn version 3.0

## ERRORED SEQUENCES

127 <210> SEQ ID NO: 10  
 128 <211> LENGTH: 25  
 129 <212> TYPE: DNA  
 C--> 130 <213> ORGANISM: Artificial<220>  
 W--> 131 <220> FEATURE:  
 131 <223> OTHER INFORMATION: Antisense Primer  
 E--> 133 <400> SEQUENCE: 10  
 134 ctttaagaag gagatgccac catgg

*Insert hard return*

25

RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 08/06/2002  
PATENT APPLICATION:    US/10/069,111A      TIME: 16:29:02

Input Set : A:\p22033.ST25.txt  
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,11,12,13,14,15,16

## VERIFICATION SUMMARY

DATE: 08/06/2002

PATENT APPLICATION: US/10/069,111A

TIME: 16:29:02

Input Set : A:\p22033.ST25.txt

Output Set: N:\CRF3\08062002\J069111A.raw

L:5 M:283 W: Missing Blank Line separator, <120> field identifier  
L:6 M:283 W: Missing Blank Line separator, <130> field identifier  
L:7 M:283 W: Missing Blank Line separator, <140> field identifier  
L:11 M:283 W: Missing Blank Line separator, <160> field identifier  
L:17 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1  
L:18 M:283 W: Missing Blank Line separator, <220> field identifier  
L:30 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2  
L:31 M:283 W: Missing Blank Line separator, <220> field identifier  
L:41 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3  
L:42 M:283 W: Missing Blank Line separator, <220> field identifier  
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:57 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4  
L:58 M:283 W: Missing Blank Line separator, <220> field identifier  
L:67 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:73 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5  
L:74 M:283 W: Missing Blank Line separator, <220> field identifier  
L:84 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6  
L:85 M:283 W: Missing Blank Line separator, <220> field identifier  
L:95 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7  
L:96 M:283 W: Missing Blank Line separator, <220> field identifier  
L:108 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8  
L:109 M:283 W: Missing Blank Line separator, <220> field identifier  
L:119 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9  
L:120 M:283 W: Missing Blank Line separator, <220> field identifier  
L:130 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10  
L:131 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:10  
L:133 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10  
L:140 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11  
L:141 M:283 W: Missing Blank Line separator, <220> field identifier  
L:151 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12  
L:152 M:283 W: Missing Blank Line separator, <220> field identifier  
L:162 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13  
L:163 M:283 W: Missing Blank Line separator, <220> field identifier  
L:173 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14  
L:174 M:283 W: Missing Blank Line separator, <220> field identifier  
L:184 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15  
L:185 M:283 W: Missing Blank Line separator, <220> field identifier  
L:195 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16  
L:196 M:283 W: Missing Blank Line separator, <220> field identifier